SEQUENCE LISTING

<110> Pan, Yang

<120> NOVEL MOLECULES OF THE TANGO-93-RELATED PROTEIN FAMILY AND USES THEREOF

<130> 07334-369001

<140> US 10/134,410
<141> 2002-04-29

<150> US 09/131,263
<151> 1998-08-07

<150> US 09/369,693
<151> 1999-08-06

<160> 14

<170> FastSEQ for Windows Version 4.0

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<212> DNA
<213> Mus musculus

<220>
<221> CDS
<222> (137)...(604)

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caaattttcc agccttgtct ttgcctaaaa tttcctgctg tttatttcaa aatagggtct 120
acatactgtg gagctc atg atg gtt ctg agt ggg gca cta tgc ttc cga atg 172
Met Met Val Leu Ser Gly Ala Leu Cys Phe Arg Met
1 5 10

aag gat tca gcc ttg aag gta ctg tat ctg cac aat aac cag ctg ctg 220
Lys Asp Ser Ala Leu Lys Val Leu Tyr Leu His Asn Asn Gln Leu Leu
15 20 25

gct gga gga ctg cac gca gag aag gtc att aaa ggt gag gag atc agt 268
Ala Gly Gly Leu His Ala Glu Lys Val Ile Lys Gly Glu Glu Ile Ser
30 35 40

gtt gtc cca aat cgg gca ctg gat gcc agt ctg tcc cct gtc atc ctg 316
Val Val Pro Asn Arg Ala Leu Asp Ala Ser Leu Ser Pro Val Ile Leu
45 50 55 60

ggc gtt caa gga gga agc cag tgc cta tct tgt ggg aca gag aaa ggg 364
Gly Val Gln Gly Gly Ser Gln Cys Leu Ser Cys Gly Thr Glu Lys Gly
65 70 75

cca att ctg aaa ctt gag cca gtg aac atc atg gag ctc tac ctc ggg 412
Pro Ile Leu Lys Leu Glu Pro Val Asn Ile Met Glu Leu Tyr Leu Gly

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            80                  85                  90

gcc aag gaa tca aag agc ttc acc ttc tac cgg cgg gat atg ggt ctt      460
Ala Lys Glu Ser Lys Ser Phe Thr Phe Tyr Arg Arg Asp Met Gly Leu
         95                  100                 105

acc tcc agc ttc gaa tcc gct gcc tac cca ggc tgg ttc ctc tgc acc      508
Thr Ser Ser Phe Glu Ser Ala Ala Tyr Pro Gly Trp Phe Leu Cys Thr
    110                 115                 120

tca ccg gaa gct gac cag cct gtc agg ctc act cag atc cct gag gac      556
Ser Pro Glu Ala Asp Gln Pro Val Arg Leu Thr Gln Ile Pro Glu Asp
125                 130                 135                 140

ccc gcc tgg gat gct ccc atc aca gac ttc tac ttt cag cag tgt gac      604
Pro Ala Trp Asp Ala Pro Ile Thr Asp Phe Tyr Phe Gln Gln Cys Asp
                145                 150                 155

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gatagaggat agagagacag aggagctcca cagtaggtgg cttactcctc tccttcccta    724
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gctggctggg ccctttccct caacctttct gacatctgca gcctctctca ttcttgcctt   1084
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acactgtcct ggtttgaaac cagaggggac aataaaaaac cctgattctg gtctctactc   1204
acataaaaag aagcttgtga acattaagtg ggaagagatt gctactaaat aacatacctt   1264
ggaatttcat cttaattaaa atatacttct ctatattata tattttaaaa aaaaaaaaaa   1324
aaaaaaaaaa aaaaaaaaaa aaaaaacatg cggccg                             1360
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<210> 2
<211> 156
<212> PRT
<213> Mus musculus

<400> 2

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Leu Lys Val Leu Tyr Leu His Asn Asn Gln Leu Leu Ala Gly Gly Leu
            20                  25                  30
His Ala Glu Lys Val Ile Lys Gly Glu Glu Ile Ser Val Val Pro Asn
        35                  40                  45
Arg Ala Leu Asp Ala Ser Leu Ser Pro Val Ile Leu Gly Val Gln Gly
    50                  55                  60
Gly Ser Gln Cys Leu Ser Cys Gly Thr Glu Lys Gly Pro Ile Leu Lys
65                  70                  75                  80
Leu Glu Pro Val Asn Ile Met Glu Leu Tyr Leu Gly Ala Lys Glu Ser
                85                  90                  95
Lys Ser Phe Thr Phe Tyr Arg Arg Asp Met Gly Leu Thr Ser Ser Phe
            100                 105                 110
Glu Ser Ala Ala Tyr Pro Gly Trp Phe Leu Cys Thr Ser Pro Glu Ala
        115                 120                 125
Asp Gln Pro Val Arg Leu Thr Gln Ile Pro Glu Asp Pro Ala Trp Asp
    130                 135                 140
Ala Pro Ile Thr Asp Phe Tyr Phe Gln Gln Cys Asp
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145 150 155

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<211> 468
<212> DNA
<213> Mus musculus

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gaggagatca gtgttgtccc aaatcgggca ctggatgcca gtctgtcccc tgtcatcctg 180
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ttcctctgca cctcaccgga agctgaccag cctgtcaggc tcactcagat ccctgaggac 420
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<212> DNA
<213> Homo sapiens

<220>
<221> CDS
<222> (57)...(521)

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Val Leu Ser Gly Ala Leu Cys Phe Arg Met Lys Asp Ser Ala Leu Lys
5 10 15

gtg ctt tat ctg cat aat aac cag ctt cta gct gga ggg ctg cat gca 155
Val Leu Tyr Leu His Asn Asn Gln Leu Leu Ala Gly Gly Leu His Ala
20 25 30

ggg aag gtc att aaa ggt gaa gag atc agc gtg gtc ccc aat cgg tgg 203
Gly Lys Val Ile Lys Gly Glu Glu Ile Ser Val Val Pro Asn Arg Trp
35 40 45

ctg gat gcc agc ctg tcc ccc gtc atc ctg ggt gtc cag ggt gga agc 251
Leu Asp Ala Ser Leu Ser Pro Val Ile Leu Gly Val Gln Gly Gly Ser
50 55 60 65

cag tgc ctg tca tgt ggg gtg ggg cag gag ccg act cta aca cta gag 299
Gln Cys Leu Ser Cys Gly Val Gly Gln Glu Pro Thr Leu Thr Leu Glu
70 75 80

cca gtg aac atc atg gag ctc tat ctt ggt gcc aag gaa tcc aag agc 347
Pro Val Asn Ile Met Glu Leu Tyr Leu Gly Ala Lys Glu Ser Lys Ser
85 90 95

ttc acc ttc tac cgg cgg gac atg ggg ctc acc tcc agc ttc gag tcg 395
Phe Thr Phe Tyr Arg Arg Asp Met Gly Leu Thr Ser Ser Phe Glu Ser

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        100                 105                 110

gct gcc tac ccg ggc tgg ttc ctg tgc acg gtg cct gaa gcc gat cag      443
Ala Ala Tyr Pro Gly Trp Phe Leu Cys Thr Val Pro Glu Ala Asp Gln
    115                 120                 125

cct gtc aga ctc acc cag ctt ccc gag aat ggt ggc tgg aat gcc ccc      491
Pro Val Arg Leu Thr Gln Leu Pro Glu Asn Gly Gly Trp Asn Ala Pro
130                 135                 140                 145

atc aca gac ttc tac ttc cag cag tgt gac tagggcaacg tgccccccag        541
Ile Thr Asp Phe Tyr Phe Gln Gln Cys Asp
                150                 155

aactccctgg gcagagccag ctcgggtgag gggtgagtgg aggagaccca tggcggacaa    601
tcactctctc tgctctcagg acccccacgt ctgacttagt gggcacctga ccactttgtc    661
ttctggttcc cagtttggat aaattctgag atttggagct cagtccacgg tcctccccca    721
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catagagtca gggatctatg gcccttggcc cagccccacc cccttccctt taatcctgcc   1021
actgtcatat gctacctttc ctatctcttc cctcatcatc ttgttgtggg catgaggagg   1081
tggtgatgtc agaagaaatg gctcgagctc agaagataaa agataagtag ggtatgctga   1141
tcctctttta aaaacccaag atacaatcaa aatcccagat gctggtctct attcccatga   1201
aaaagtgctc atgacatatt gagaagacct acttacaaag tggcatatat tgcaatttat   1261
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gc                                                                  1323

<210> 5
<211> 155
<212> PRT
<213> Homo sapiens

<400> 5
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            20                  25                  30
Ala Gly Lys Val Ile Lys Gly Glu Glu Ile Ser Val Val Pro Asn Arg
        35                  40                  45
Trp Leu Asp Ala Ser Leu Ser Pro Val Ile Leu Gly Val Gln Gly Gly
    50                  55                  60
Ser Gln Cys Leu Ser Cys Gly Val Gly Gln Glu Pro Thr Leu Thr Leu
65                  70                  75                  80
Glu Pro Val Asn Ile Met Glu Leu Tyr Leu Gly Ala Lys Glu Ser Lys
                85                  90                  95
Ser Phe Thr Phe Tyr Arg Arg Asp Met Gly Leu Thr Ser Ser Phe Glu
            100                 105                 110
Ser Ala Ala Tyr Pro Gly Trp Phe Leu Cys Thr Val Pro Glu Ala Asp
        115                 120                 125
Gln Pro Val Arg Leu Thr Gln Leu Pro Glu Asn Gly Gly Trp Asn Ala
    130                 135                 140
Pro Ile Thr Asp Phe Tyr Phe Gln Gln Cys Asp
145                 150                 155

<210> 6
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<211> 465
<212> DNA
<213> Homo sapiens

<400> 6
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gagatcagcg tggtccccaa tcggtggctg gatgccagcc tgtcccccgt catcctgggt 180
gtccagggtg gaagccagtg cctgtcatgt ggggtggggc aggagccgac tctaacacta 240
gagccagtga acatcatgga gctctatctt ggtgccaagg aatccaagag cttcaccttc 300
taccggcggg acatggggct cacctccagc ttcgagtcgg ctgcctaccc gggctggttc 360
ctgtgcacgg tgcctgaagc cgatcagcct gtcagactca cccagcttcc cgagaatggt 420
ggctggaatg cccccatcac agacttctac ttccagcagt gtgac 465

<210> 7
<211> 177
<212> PRT
<213> Homo sapiens

<400> 7
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Phe Leu Phe His Ser Glu Thr Ile Cys Arg Pro Ser Gly Arg Lys Ser
20 25 30
Ser Lys Met Gln Ala Phe Arg Ile Trp Asp Val Asn Gln Lys Thr Phe
35 40 45
Tyr Leu Arg Asn Asn Gln Leu Val Ala Gly Tyr Leu Gln Gly Pro Asn
50 55 60
Val Asn Leu Glu Glu Lys Ile Asp Val Val Pro Ile Glu Pro His Ala
65 70 75 80
Leu Phe Leu Gly Ile His Gly Gly Lys Met Cys Leu Ser Cys Val Lys
85 90 95
Ser Gly Asp Glu Thr Arg Leu Gln Leu Glu Ala Val Asn Ile Thr Asp
100 105 110
Leu Ser Glu Asn Arg Lys Gln Asp Lys Arg Phe Ala Phe Ile Arg Ser
115 120 125
Asp Ser Gly Pro Thr Thr Ser Phe Glu Ser Ala Ala Cys Pro Gly Trp
130 135 140
Phe Leu Cys Thr Ala Met Glu Ala Asp Gln Pro Val Ser Leu Thr Asn
145 150 155 160
Met Pro Asp Glu Gly Val Met Val Thr Lys Phe Tyr Phe Gln Glu Asp
165 170 175
Glu

<210> 8
<211> 178
<212> PRT
<213> Mus musculus

<400> 8
Met Glu Ile Cys Trp Gly Pro Tyr Ser His Leu Ile Ser Leu Leu Leu
1 5 10 15
Ile Leu Leu Phe His Ser Glu Ala Ala Cys Arg Pro Ser Gly Lys Arg
20 25 30
Pro Cys Lys Met Gln Ala Phe Arg Ile Trp Asp Thr Asn Gln Lys Thr
35 40 45

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Phe Tyr Leu Arg Asn Asn Gln Leu Ile Ala Gly Tyr Leu Gln Gly Pro
    50                  55                  60
Asn Ile Lys Leu Glu Glu Lys Ile Asp Met Val Pro Ile Asp Leu His
65                  70                  75                  80
Ser Val Phe Leu Gly Ile His Gly Gly Lys Leu Cys Leu Ser Cys Ala
                85                  90                  95
Lys Ser Gly Asp Asp Ile Lys Leu Gln Leu Glu Glu Val Asn Ile Thr
            100                 105                 110
Asp Leu Ser Lys Asn Lys Glu Glu Asp Lys Arg Phe Thr Phe Ile Arg
        115                 120                 125
Ser Glu Lys Gly Pro Thr Thr Ser Phe Glu Ser Ala Ala Cys Pro Gly
    130                 135                 140
Trp Phe Leu Cys Thr Thr Leu Glu Ala Asp Arg Pro Val Ser Leu Thr
145                 150                 155                 160
Asn Thr Pro Glu Glu Pro Leu Ile Val Thr Lys Phe Tyr Phe Gln Glu
                165                 170                 175
Asp Gln
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<210> 9
<211> 13
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetically generated primer

<400> 9
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<210> 10
<211> 17
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetically generated primer

<400> 10
cacctcgagt actaccc 17

<210> 11
<211> 16
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetically generated primer

<400> 11
cgaggtctac caggac 16

<210> 12
<211> 16
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetically generated primer

<400> 12
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<210> 13
<211> 2490
<212> DNA
<213> Homo sapiens

<400> 13
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atctgcataa taaccagctt ctagctggag ggctgcatgc agggaaggtc attaaaggtg 180
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<210> 14
<211> 155

<212> PRT
<213> Homo sapiens

<400> 14

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            20                  25                  30
Ala Gly Lys Val Ile Lys Gly Glu Glu Ile Ser Val Val Pro Asn Arg
        35                  40                  45
Trp Leu Asp Ala Ser Leu Ser Pro Val Ile Leu Gly Val Gln Gly Gly
    50                  55                  60
Ser Gln Cys Leu Ser Cys Gly Val Gly Gln Glu Pro Thr Leu Thr Leu
65                  70                  75                  80
Glu Pro Val Asn Ile Met Glu Leu Tyr Leu Gly Ala Lys Glu Ser Lys
                85                  90                  95
Ser Phe Thr Phe Tyr Arg Arg Asp Met Gly Leu Thr Ser Ser Phe Glu
            100                 105                 110
Ser Ala Ala Tyr Pro Gly Trp Phe Leu Cys Thr Val Pro Glu Ala Asp
        115                 120                 125
Gln Pro Val Arg Leu Thr Gln Leu Pro Glu Asn Gly Gly Trp Asn Ala
    130                 135                 140
Pro Ile Thr Asp Phe Tyr Phe Gln Gln Cys Asp
145                 150                 155
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